> This Raw Listing contains the General Information
> Section and up to first 5 pages.

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  1   <110> APPLICANT: Xu, Jiangchun
  2           Dillon, Davin C.
  3           Mitcham, Jennifer Lynn
  4   <120> TITLE OF INVENTION: COMPOUNDS FOR IMMUNODIAGNOSIS OF
  5           PROSTATE CANCER AND METHODS FOR THEIR USE
  6   <130> FILE REFERENCE: 210121.428C6
  7   <140> CURRENT APPLICATION NUMBER: US/09/232,880
  8   <141> CURRENT FILING DATE: 1999-01-15
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 32         actcctcaaa ggnggtatta cggttatccn naaatcnggg gatacccngg aaaaaanttt     780
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ENTERED

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PREVIOUSLY ERRORED SEQUENCES-EDITED

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23       tccaaccntc gntggccntn ccccccncnn tcctttnccc                          820
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CRF E☰☰'s Corrected by the STIC System ☰ranch

Serial Number: _09/232,880_

C☰☰ Processing Date: _4/22/99_
Edited by: _____
Verified by: _____ (STIC staff)

☐ Changed a file from non-ASCII to ASCII

☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.

☐ Edited a format error in the Current Application Data section, specifically:
_____

☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____

~~ENTERED~~

☐ Added the mandatory heading and subheadings for "Current Application Data".

☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.

☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically:
_____

☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:
_____

☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:
_____

☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.

☐ Inserted colons after headings/subheadings. Headings edited included:
_____

☐ Deleted extra, invalid, headings used by an applicant, specifically:
_____

☐ Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as_____ .

☐ Inserted mandatory headings, specifically: _____

☐ Corrected an obvious error in the response, specifically:
_____

☐ Edited identifiers where upper case is used but lower case is required, or vice versa.

☐ Corrected an error in the Number of Sequences field, specifically:
_____

☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.

☐ Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____

☑ Other: _pg 26 - inserted hard return after cumulative loss total 780⁵_
_____
_____

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.   3/1/95

Input Set: **I232880.RAW**

> **This Raw Listing contains the General Information Section and those Sequences containing ERRORS.**

Corrected Does Not Comply Diskette Needed

1	<110>	Xu, Jiangchun
2		Dillon, Davin C.
3		Mitcham, Jennifer Lynn
4	<120>	COMPOUNDS FOR IMMUNODIAGNOSIS OF
5		PROSTATE CANCER AND METHODS FOR THEIR USE
6	<130>	210121.428C6
7	<140>	US/09/232,880
8	<141>	1999-01-15
9	<160>	338
10	<170>	FastSEQ for Windows Version 3.0

ERRORED SEQUENCES FOLLOW

	11	<210>	26
	12	<211>	820
	13	<212>	DNA
	14	<213>	Homo sapien
	15	<220>	
	16	<221>	misc_feature
	17	<222>	(1)...(820)
	18	<223>	n = A,T,C or G
	19	<400>	26
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W-->	21	cccanagata ncttatanca acagtgcttt gaccaagagc tgctgggcac atttcctgca	120
	22	gaaaaggtgg cggtccccat cactcctcct ctcccatagc catcccagag gggtgagtag	180
W-->	23	ccatcangcc ttcggtggga gggagtcang gaaacaacan accacagagc anacagacca	240
W-->	24	ntgatgacca tgggcgggag cgagcctctt ccctgnaccg gggtggcana nganagccta	300
W-->	25	nctgaggggt cacactataa acgttaacga ccnagatnan cacctgcttc aagtgcaccc	360
W-->	26	ttcctacctg acnaccagng accnnnaact gcngcctggg gacagcnctg ggancagcta	420
W-->	27	acnnagcact cacctgcccc cccatggccg tncgcntccc tggtcctgnc aagggaagct	480
W-->	28	ccctgttgga attncgggga naccaaggga nccccctcct ccanctgtga aggaaaaann	540
W-->	29	gatggaattt tnccttccg gccnntcccc tcttccttta cacgcccct nntactcntc	600
W-->	30	tccctctntt ntcctgncnc acttttnacc ccnnnatttc ccttnattga tcggannctn	660
W-->	31	ganattccac tnncgcctnc cntcnatcng naanacnaaa nactntctna cccnggggat	720
E-->	32	gggnncctcg ntcatcctct ctttttcnct accnccnntt ctttgcctct ccttngatca	780tcca

Line	?	Error/Warning	Original Text
20	W	"N" or "Xaa" used: Feature required	anattantac agtgtaatct tttcccagag gtgtgtan
21	W	"N" or "Xaa" used: Feature required	cccanagata ncttatanca acagtgcttt gaccaaga
23	W	"N" or "Xaa" used: Feature required	ccatcangcc ttcggtggga gggagtcang gaaacaac
24	W	"N" or "Xaa" used: Feature required	ntgatgacca tgggcgggag cgagcctctt ccctgnac
25	W	"N" or "Xaa" used: Feature required	nctgaggggt cacactataa acgttaacga ccnagatn
26	W	"N" or "Xaa" used: Feature required	ttcctacctg acnaccagng accnnnaact gcngcctg
27	W	"N" or "Xaa" used: Feature required	acnnagcact cacctgcccc cccatggccg tncgcntc
28	W	"N" or "Xaa" used: Feature required	ccctgttgga attncgggga naccaaggga nccccctc
29	W	"N" or "Xaa" used: Feature required	gatggaattt tncccttccg gccnntcccc tcttcctt
30	W	"N" or "Xaa" used: Feature required	tccctctntt ntcctgncnc acttttnacc ccnnnatt
31	W	"N" or "Xaa" used: Feature required	ganattccac tnncgcctnc cntcnatcng naanacna
32	E	Number of Bases conflict w/ Running Total	gggnncctcg ntcatcctct ctttttcnct accnccnn
32	E	Wrong Nucleic Acid Designator	gggnncctcg ntcatcctct ctttttcnct accnccnn
32	E	Wrong Nucleic Acid Designator	gggnncctcg ntcatcctct ctttttcnct accnccnn
32	E	Wrong Nucleic Acid Designator	gggnncctcg ntcatcctct ctttttcnct accnccnn
32	W	"N" or "Xaa" used: Feature required	gggnncctcg ntcatcctct ctttttcnct accnccnn